101 IVRGTQLFED NYALAVLDNG DPLNNTTPVT GASPGGLREL QLRSLTEILK GGVLIQRNPQ LCYQDTILWK DIFHKNNQLA LTLIDTNRSR ACHPCSPMCK 401 ETLEEITGYL YISAWPDSLP DLSVFQNLQV IRGRILHNGA YSLTLQGLGI SWLGLRSLRE LGSGLALIHH NTHLCFVHTV PWDQLFRNPH QALLHTANRP 301 YNYLSTDVGS CTLVCPLHNQ EVTAEDGTQR CEKCSKPCAR VCYGLGMEHL REVRAVTSAN IQEFAGCKKI FGSLAFLPES FDGDPASNTA PLQPEQLQVF 201 GSRCWGESSE DCQSLTRTVC AGGCARCKGP LPTDCCHEQC AAGCTGPKHS DCLACLHFNH SGICELHCPA LVTYNTDTFE SMPNPEGRYT FGASCVTACP 601 PSGVKPDLSY MPIWKFPDEE GACQPCPINC THSCVDLDDK GCPAE (SEG NO NO: N3) 501 EDECVGEGLA CHQLCARGHC WGPGPTQCVN CSQFLRGQEC VEECRVLQGL PREYVNARHC LPCHPECQPQ NGSVTCFGPE ADQCVACAHY KDPPFCVARC 1 MELAALCRWG LLLALLPPGA ASTQVCTGTD MKLRLPASPE THLDMLRHLY QGCQVVQGNL ELTYLPTNAS LSFLQDIQEV QGYVLIAHNQ VRQVPLQRLR

ig. 1A

(31 residues) 7C2 aa 22-53 aa 22-53 (31 residues) 7F3 (562 residues) 2C4 aa 22-584 (562 residues) 7D3 aa 22-584 aa 512-625 (113 residues) 3E8 (96 residues) 4D5 aa 529-625 2H11 aa 529-645 (116 residues) (58 residues) 3H4 aa 541-599

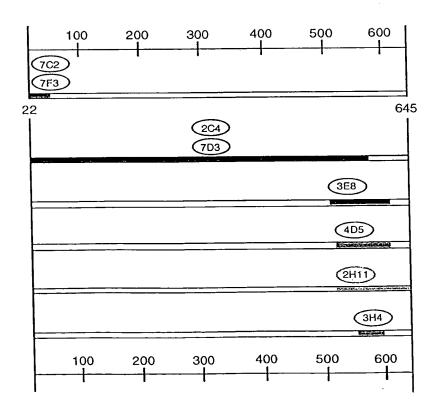


Fig. 1B

CTECHOLD CENTUCE

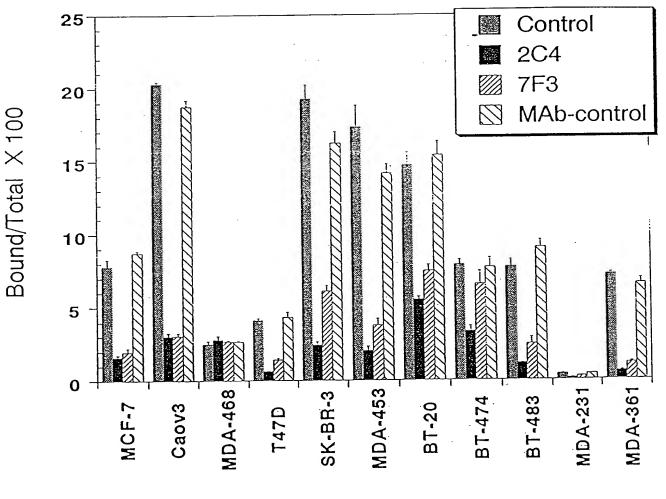
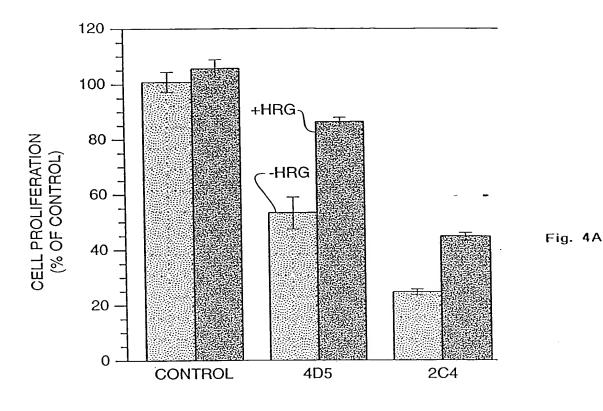
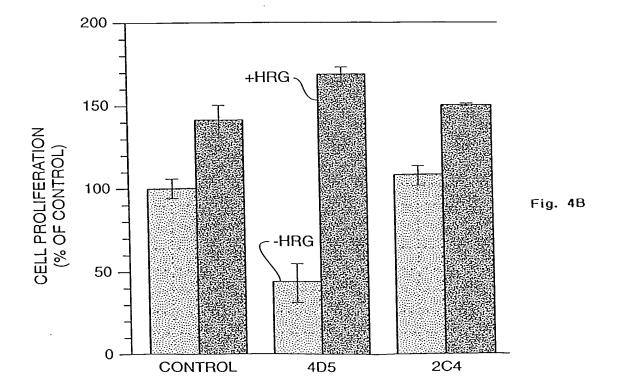


Fig. 3



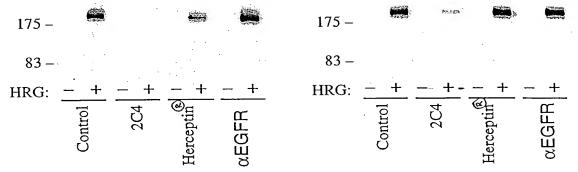


MCF7
Low/Normal HER2

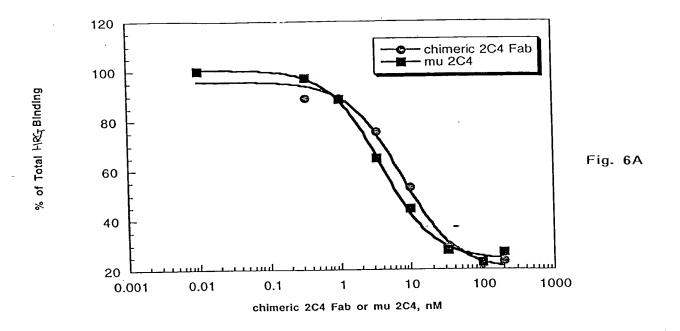
Fig. 5A

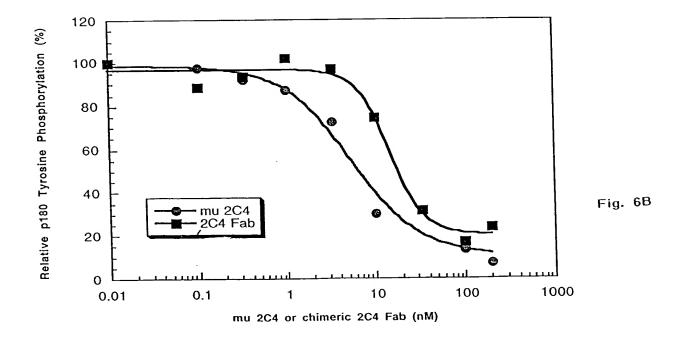
SK-BR-3 High HER2

Fig. 5B



IP: αHER2 WB: αHER3





hum III

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Fig. 7A: Variable Light
                           20
                                        30
                                                    40
                 10
         DTVMTOSHKIMSTSVGDRVSITC [KASQDVSIGVA] WYQQRP
2C4
                ****
         DIQMTQSPSSLSASVGDRVTITC [KASQDVSIGVA] WYQQKP
574
         DIOMTQSPSSLSASVGDRVTITC [RASQSISNYLA] WYQQKP
hum KI
                                                    80
                   50
                                60
                                          70
         GQSPKLLIY [SASYRYT] GVPDRFTGSGSGTDFTFTISSVQA
2C4
         GKAPKLLIY [SASYRYT] GVPSRFSGSGSGTDFTLTISSLQP
574
         GKAPKLLIY [AASSLES] GVPSRFSGSGSGTDFTLTISSLQP
hum KI
                               100
                    90
         EDLAVYYC [QQYYIYPYT] FGGGTKLEIK (SEQ ID NO:1)
2C4
         EDFATYYC [QQYYIYPYT] FGQGTKVEIK (SEQ ID NO:3)
574
         EDFATYYC [QQYNSLPWT] FGQGTKVEIK (SEQ ID NO:5)
hum ĸI
Fig. 7B: Variable Heavy
                            20
                                         30
                  10
          EVOLOQSGPELVKPGTSVKISCKAS [GFTFTDYTMD] WVKQS
2C4
                 ** * * ***
          EVQLVESGGGLVQPGGSLRLSCAAS [GFTFTDYTMD] WVRQA
574
         EVOLVESGGGLVQPGGSLRLSCAAS [GFTFSSYAMS] WVRQA
hum III
                                            70
                    50
                               60
                       а
          HGKSLEWIG [DVNPNSGGSIYNQRFKG] KASLTVDRSSRIVYM
 2C4
          PGKGLEWVA [DVNPNSGGSIYNQRFKG] RFTLSVDRSKNTLYL
 574
                     *****
                                            * * *
          PGKGLEWVA [VISGDGGSTYYADSVKG] RFTISRDNSKNTLYL
 hum III
                                               110
            abc
                     90
                                 100ab
          ELRSLTFEDTAVYYCAR [NLGPSFYFDY] WGQGTTLTVSS (SEQ ID NO:2)
 2C4
                                               * *
          QMNSLRAEDTAVYYCAR [NLGPSFYFDY] WGQGTLVTVSS (SEQ ID NO:4)
 574
          QMNSLRAEDTAVYYCAR [GRVGYSLYDY] WGQGTLVTVSS (SEQ ID NO:6)
```

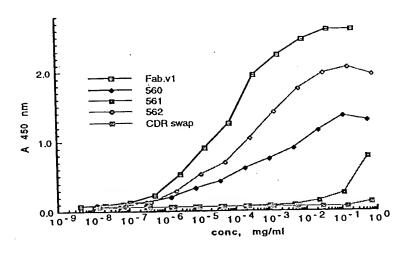


Fig. 8A

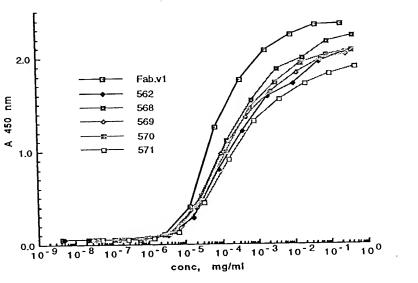
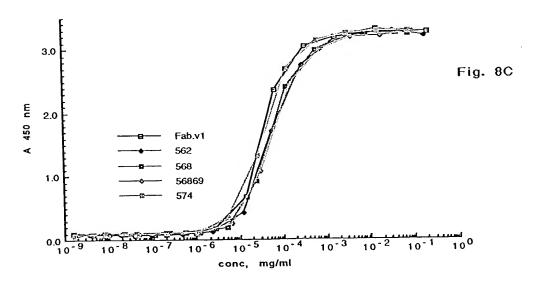


Fig. 8B



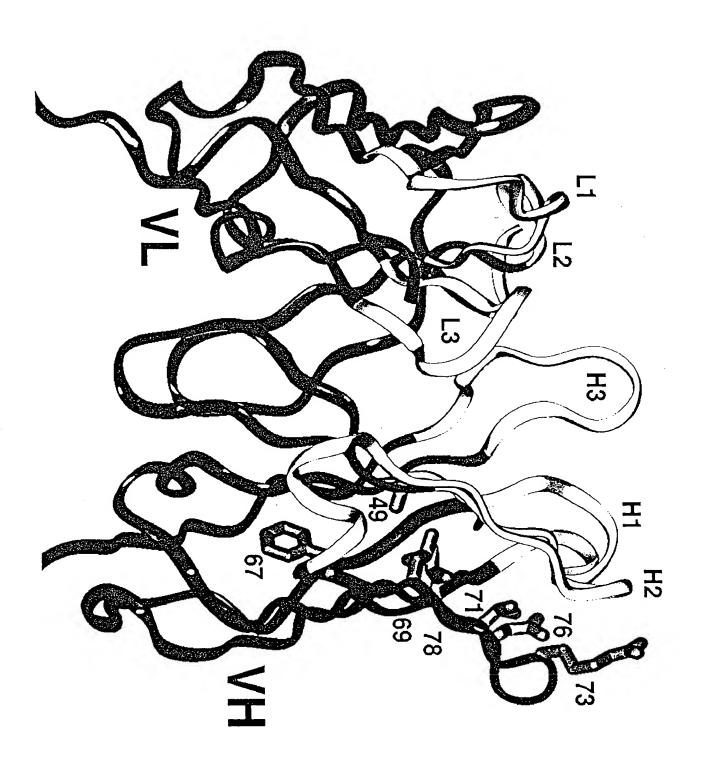
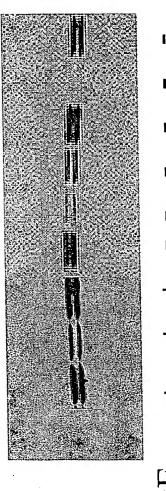
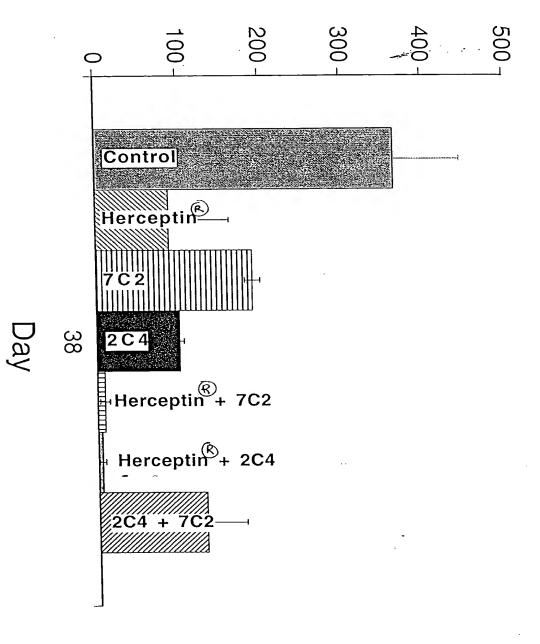


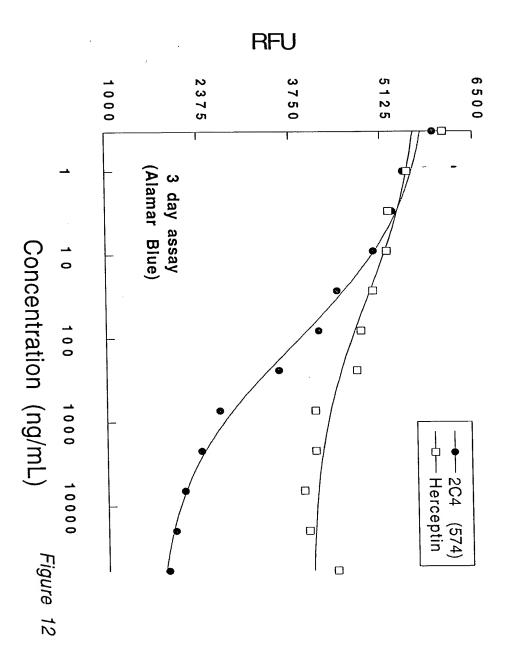
Fig. 10



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	I	+	1	1	i
	+	1	ı	1	+
	+	ſ	ï	+	I
	+	ı	ı	i	I
Marija da	1	ı	ı	ı	ı
	EGF	$TGF\alpha$	HRG	2C4	FRCEPTING



HIG. II



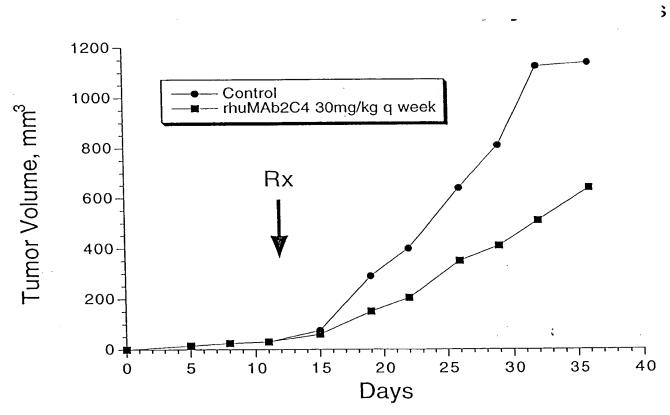


Fig. 13